## Twenty Years of High Performance, Parallel Computing: Vector Machines to HPC Linux Clusters

#### **Harold Trease**

Computation Science and Mathematics Division Fundamental Sciences Directorate Pacific Northwest National Laboratory





#### **Overview**

- ► This is "not" meant to be a stroll down memory lane.
- ► Hopefully, I can convey some useful lessons learned.
- I work at the application software level.
- ► I try to influence the development of "middleware".
- I have to live with the delivered hardware.
- I want to do application software development using a "shared memory" programming model, but have access to high-performance parallel computing resources for solving application problems. And I won't be happy until I can do this at the petabytes/petaflop scale.

## Why do this level of HPC computing?

- Scientific Discovery through Advanced Computing
  - Science drivers
  - Multi-scale (time and/or space) resolution
  - Complex physics
- Computational science reasons:
  - Benchmarking
  - Turnaround time
  - Access of large memory (←)
  - Cycle sucking parasite



## **Overview**

- Review of the hardware
- Software of the software
  - Codes
  - Languages
  - Parallel Programming Paradigms
- **►** Environments
- ► Soapbox Issues
- Applications
- Lessons Learned



## Hardware Background (8 strong oxen .vs. 10,000 chickens)

#### LANL:

- CRAY Series [1s, XMP, YMP] at LANL (PCs days) [80's]
- Connection Machines [CM2/CM5] [late 80's early 90's]
- Cray T3D/T3E series [early middle 90's]
- SGI ASCI Blue Mountain [middle late 90's]

#### ► PNNL:

- IBM SP2 [early 2000's]
- PC Cluster Computing [early 2000's]
- ORNL IBM/Power4 and PNNL HP (current)



## **Hardware Background**

- CRAYs: Evolving, stable software development environment, 64-bit data types, gather-scatter, multi-tasking
- ► CM2: Data Parallel, SIMD, RTS, F90, Global operations
- ► CM5 [MIMD, MPI]
- ► T3D/T3E [shmem]
- ➤ SGI Blue Mountain [Terabytes/Teraflops, dedicated time for terascale application runs, billion element meshes]
- ► IBM SP2 [32-bits]
- PC Cluster Computing [Price/Performance]
- ► HP Super Cluster [TBD, but 10s of Terabytes/Teraflops]



## Software Background

- Computational mesh based methods:
  - Unstructured 3-D meshes, compressed sparse matrix data structures (unassembled/assembled), dynamic in space and time.
- Computational geometry
  - → Computational meshes
    - → Computational physics solvers
- One semi-continuous computational physics code development effort.
  - FLM2D → FLM3D→ POLLY→ X3D→ NWGrid→ P3D
  - ~400 man-years, 2-3 million lines of core code
  - 40-50 developers have added/contributed code



## Languages

► 80's: F77, CRAY style pointers, dynamic memory allocation, runtime database management support

▶ 90's: CMFortran, C++(BS), F90, Python

► 2000's: Components using a mixture of languages (F77/F90, C, C++, Python, Java) through the Common Component Architecture

## **Programming Paradigms**

- Multiple vector processes that parallel process through time-slicing and communicate through disk files. [CRAY-1]
- Vector/Multitasking [CRAY-XMP/YMP]
- Data Parallel, SIMD [CM2]
- ► MIMD, MPI [CM5, T3D/E] (the dark times)
- Data Parallel, MIMD [SGI, HP]

## (My) Programming Paradigm

- Domain decomposition (partitioning)
  - Load balancing
  - Maximum locality of data
- Data parallel processor-to-processor communication through a library interface called Global Arrays.
- Language
  - F77/F90, C, C++
  - "Embedded" Python with shared memory access
  - Runtime, memory resident relational database (RDBMS)
- ► Components (CCA)

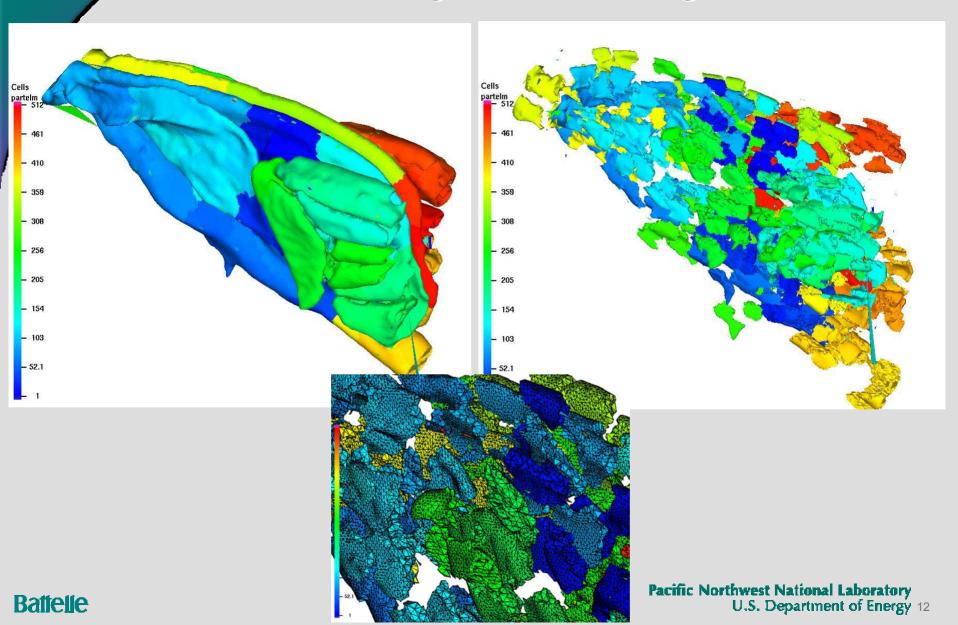


# Global Arrays: A data parallel communication library for managing distributed shared memory data structures.

- Gets/Puts, Gather/Scatter, etc.
- ► Non-uniform data layouts
- Scatter\_with\_sum
- Segmented Scans (operators=sum, copy, max/min)
- Compress/Uncompress
- Two key sorter (e.g., used to map from nodes/elements to rows and columns of a compressed sparse matrix)
- M X N mapping



## Maximize Locality of Data using METIS



### **Environments**

- Operating Systems, Schedulers, Batch Management:
  - Important, but I don't have much impact on the choice?
  - Always report security issues, but never report other anomalies because they might come in handy.
- ► I/O Issues: I prefer global shared disks, but don't require them.
- Compilers: Pretty good for what they do. They only serve the purpose of giving me access to my RDBMS.



## **Environments (cont)**

- Multi-processor, Dynamic Debugging Environment:
  - Why can't we have one available before a machine reaches the end of it's lifetime ??
  - Checkpoint/Restart (available on early CRAYs)
- Distributed Graphics/Visualization: Solutions get tough for terascale datasets (like billion elements meshes and mesh data).



# Computational Infrastructure and Applications

► DOE Office of Science

►PNNL EMSL

► HPCS2 and PNNL Computing Infrastructure



#### The DOE Office of Science has three centers with distinct architectural features to enable scientific discovery through advanced computing

#### **EMSL-HP**

#### 11 teraflops



- ➤ HP Itanium2
- ► 1,980 processors
- ► 6 GFLOP processor
- ▶ 98% DGEMM efficiency
- ► 125 cabinets

#### **NERSC-IBM**

#### 10 teraflops



- ► IBM Power3
- ► 6,656 processors
- ► 1.5 GFLOP processor
- ► 87% DGEMM efficiency
- ► 104 cabinets

#### **ORNL-Cray/IBM**

#### 3.2 teraflops



4.5 teraflops



Cray X1

- ► IBM Power4
- ► 256 processors
- ► 864 processors
- ► 12.8 GFLOP processor ► 5.2 GFLOP processor
- ➤ ?? DGEMM efficiency ➤ 65% DGEMM efficiency

8 cabinets

30 cabinets

#### **Three Centers - Three Vendors**

## **EMSL Lab Instruments**



**High Field Magnetic Resonance Facility** 

**Chemistry and Physics of Complex Systems Facility** 

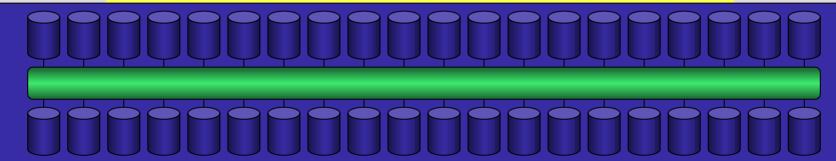


Interfacial and Nanoscale **Science Facility** 





**High Performance Mass Spectrometry Facility** 

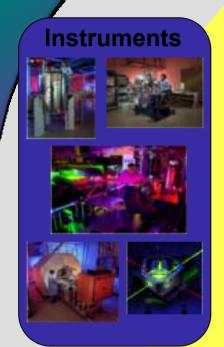


**END-TO-END GIGABIT ETHERNET TCP/IP NETWORK** 

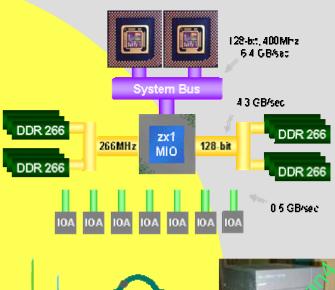
**Environmental Spectroscopy and** 

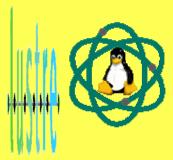
**Biogeochemistry Facility** 

## HPCS2



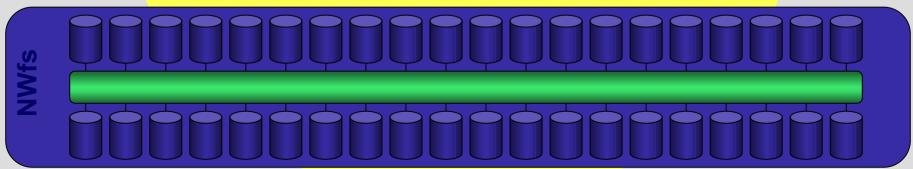
1900 Madison Processors
NWLinux
Quadric's Elan4
hp's ZX1 chip set
Gig-E connection to all nodes
53 TB of global storage
200 TG local disk
Lustre Lite File System





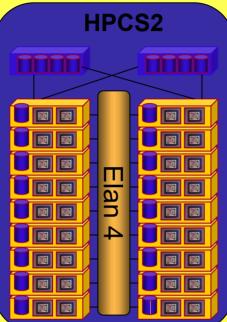


**END-TO-END GIGABIT ETHERNET TCP/IP NETWORK** 



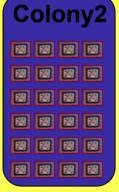
### The Grid





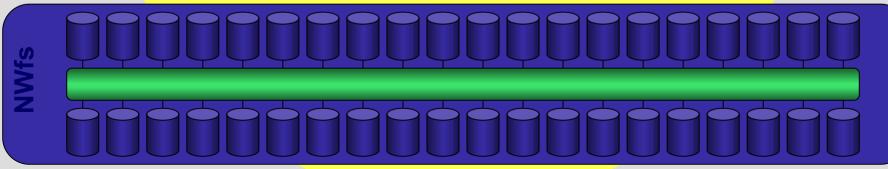
DOE Science Grid

**Internal Globus Grid** 





**END-TO-END GIGABIT ETHERNET TCP/IP NETWORK** 

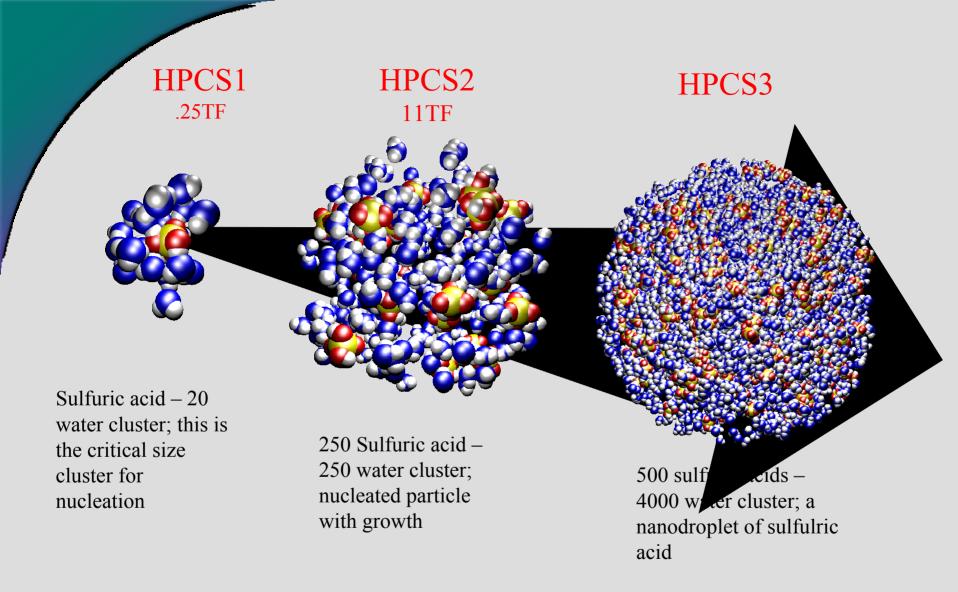




### **DOE/PNNL Science Drivers**

- ▶ Chemistry
- Subsurface Transport
- ► Atmospheric Transport
- Computational Biology

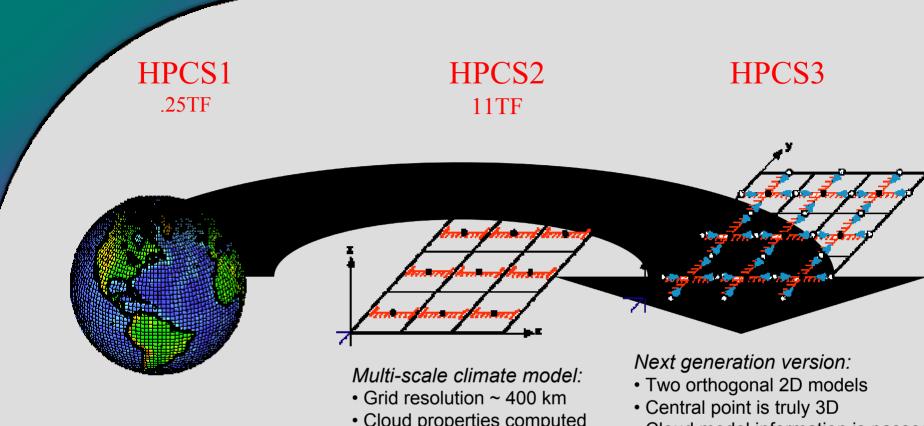




## Chemical Dynamics: (Greg Shenter)

Nucleation for aerosols





"Traditional model"

- Coarse grid resolution
- Cloud parameterization averaged over entire box

- Cloud properties computed explicitly on embedded 2D model at scale of 4 km
- Computational burden increases by factor of 200
- Can be run for 1 or 2 years of simulated time
- Cloud model information is passed across outer grid boundaries at end of time step
- Computational burden will be more than double previous model

Atmospheric Chemistry and Modeling: (Tom Ackerman)

Multi-scale climate models with increased fidelity



## HPCS1 .25TF

Subsurface simulations approaching 2 orders of magnitude of resolution in each spatial dimension



HPCS3

Complex, coupled process simulations approaching 3 orders of magnitude of resolution in each spatial direction

Complex, coupled process simulations over 2 orders of magnitude of resolution in each spatial direction

#### Contaminant behavior

Fundamental scientific basis for field-scale predictions

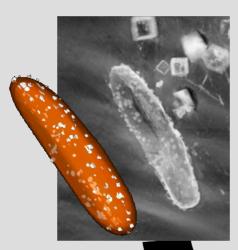


HPCS1 25TF

HPCS2 **11TF** 

HPCS3

We couldn't compute spatial details on MPP1, could generate basic geometry



Begin to mo atial details on an In cell basis with "sinbiochemistry.

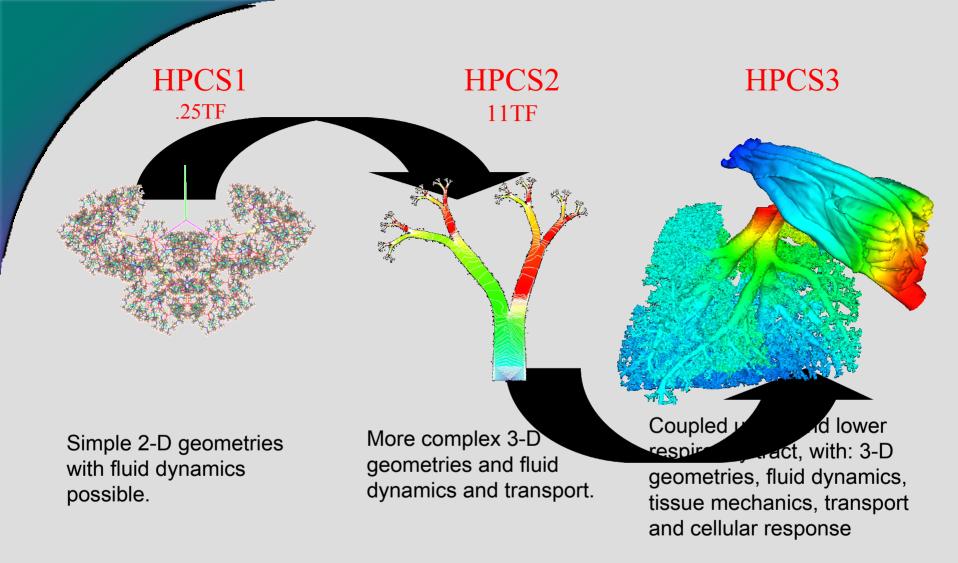


del multi-cell Potential # with "complex" commu ary within an Jonmental context.

## Computational Biology: (Harold Trease)

Modeling and Simulation of the Bioremediation of Heavy Metal Waste by Shewanella MR-1 using Multi-Scale Computational Methods



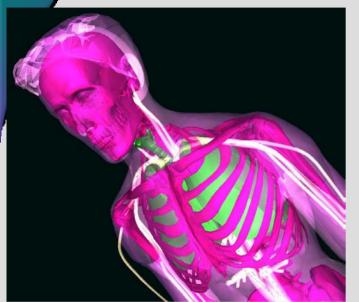


## Computational Biology: (Harold Trease)

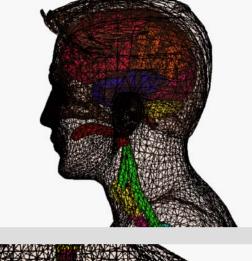
Virtual Respiratory Tract Modeling and Simulation with Particle Deposition and Chemical Vapor Transport

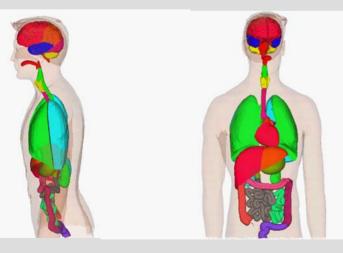


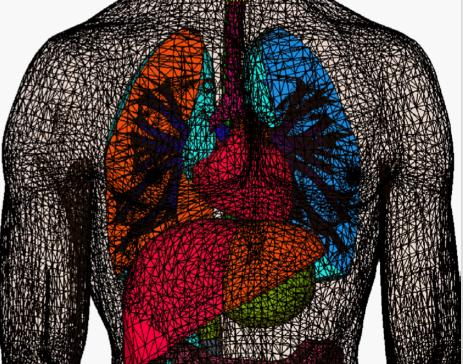
## Modeling and Simulation of Complex Biological Systems



(Frankenstein meets Extreme Computing)



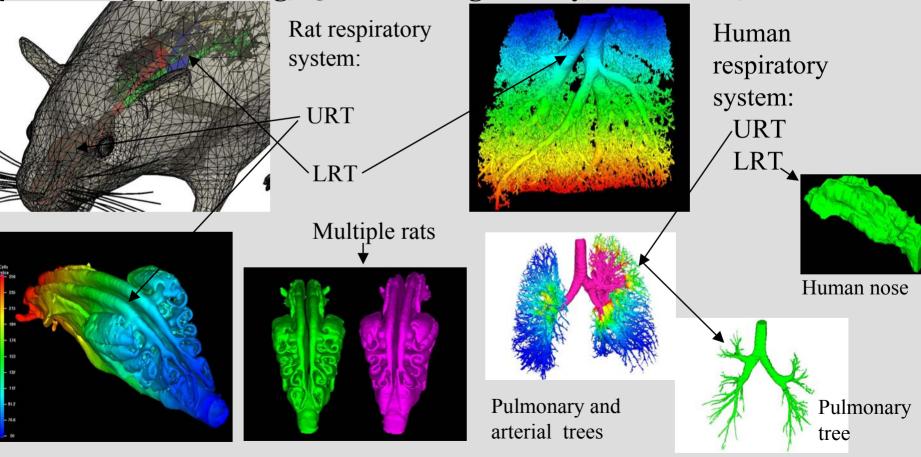




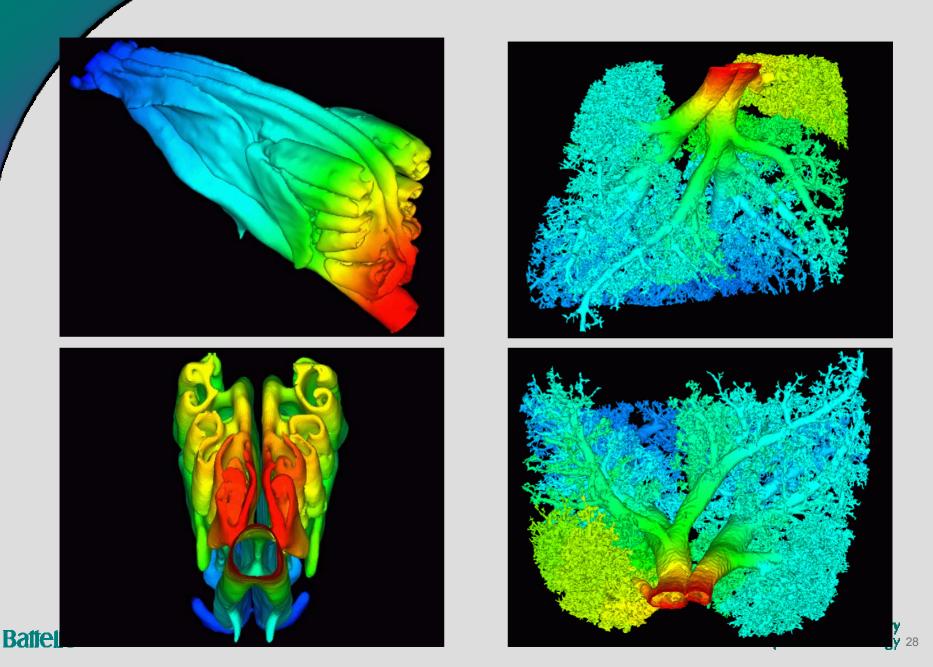
## The Virtual Respiratory Tract (VRT)

- The team: Rick Corley, Chuck Timchalk, Kevin Minard, Harold Trease, Lynn Trease)
  - Multiple Species, Multiple individuals/species

- Image processing, Quantitative geometry, Simulation, Validation



## Reconstructions of a rats nose and lungs from NMR data.



## Soapbox Issues

- ➤ We need a high-level parallel language.
  - Locality of data (may be a RTS issue)
  - Communication libraries
  - Language
- ► Dynamic Debugging Environment:
  - Why can't we have one available before a machine reaches the end of it's lifetime ??

## Soapbox Issues

► Time to solution as a metric for success. Better mathematics and numerics don't always lead to increased machine performance and higher efficiency.

Scalability: Approximate linear scaling of mesh based methods, with number of processors and problem size, is guaranteed unless you screw it up.

## Soapbox Issues

- ▶ If the goal is to use terascale machines to solve science problems, then application programmers need access to development cycles and dedicated terascale cycles.
- ➤ For sustained performance and efficiency we need a stable code development environment. [BUT: If you can't run with the big dogs you better stay on the porch.]



## **Lessons Learned**

- Scientific Discovery Through Advanced Computing
- ► Time to solution as a metric for success.

- ► The shared memory parallel programming paradigm is the correct model for mesh based computing.
- Parallel languages (we need one)
- Debugging environments / distributed graphics/visualization
- Access to xxxxflop/xxxxbyte resources.



## PNNL/DOE Funding and Acknowledgements

- SciDAC TSTT Center (Terascale Simulation Tools and Technology) [DOE OASCR/MICS]
- Microbial Cell Biology [DOE OBER/GTL], Computational Biology Simulation Frameworks [DOE OASCR/MICS]
- Virtual Respiratory Tract Project [PNNL/LDRD, NIH]